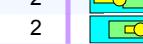
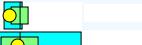
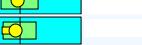
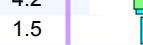
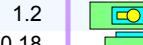
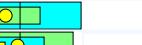
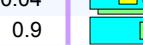
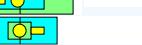
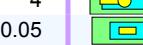
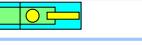
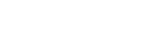


## INPUTs 2013.1

Survey	INPUTs 2013.1
Start date	January, 1 2013
Supervisor	dr. C. Weykamp MCA Laboratory Streekziekenhuis Koningin Beatrix Winterswijk
Subscriptions	93
Result sets	91

Scores	Your score	MAP	reported
Quantitative	28	38	

## INPUTs 2013.1

Analyte		Trueness				Precision		Performance				P1
		your mean	ref.	cons.	SDtl	your prec.	SDbl	this survey	PS	cumulative	PSc	
ALAT	U/l	84.3	98.2	86.9	10.1	2.3	2.7		1			1
Alk. Phosphatase	U/l	154	177	163	15	2	2		2			2
Amylase	U/l	199	207	201	8	7	2		2			2
ASAT	U/l	67.5	78.6	71.5	6.8	2.0	1.6		0			0
Calcium	mmol/l	2.45	2.46	2.47	0.07	0.06	0.02		2			2
Chloride	mmol/l	99.7	100.0	100.8	1.8	0.3	0.9		2			2
CK	U/l	230	244	253	18	4	3		2			2
Creatinine	µmol/l	157.5	149.1	150.9	7.1	7.5	4.2		0			0
eGFR (F, 55, white)	ml/min/1,73m <sup>2</sup>	50.5	37.6	37.6	2.7	2.6	1.5		0			0
Gamma-GT	U/l	82.8	87.7	86.3	8.6	2.3	1.2		2			2
Glucose	mmol/l	13.93	13.93	14.19	0.44	0.22	0.18		2			2
LD	U/l	1084	549	539	33	6	9		0			0
Magnesium	mmol/l	1.18	1.22	1.23	0.04	0.03	0.02		2			2
Potassium	mmol/l	5.38	5.39	5.30	0.10	0.03	0.04		2			2
Sodium	mmol/l	144.0	144.0	142.7	1.7	0.6	0.9		2			2
Total Protein	g/l	63.0	66.1	65.3	1.8	0.3	0.7		2			2
Urate	µmol/l	354	376	374	15	8	4		2			2
Cholesterol	mmol/l	5.07	5.07	5.15	0.14	0.06	0.05		2			2
HDL-Cholesterol	mmol/l	1.18	1.14	1.17	0.06	0.05	0.03		1			1

Total :

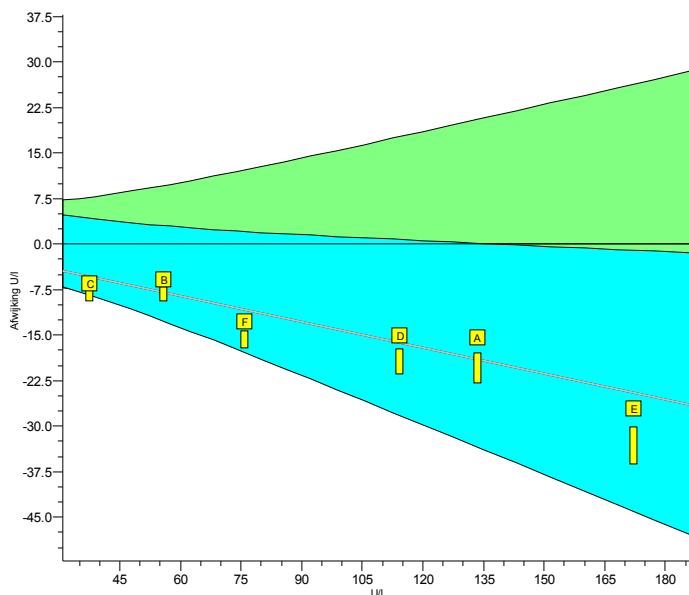
28

28

## INPUTS 2013.1

ALAT

units: U/l



	2013.1	cumulative
Trueness	-14%	-14%
Precision	2.7%	2.7%
Number	6	6
Outliers	0	0
Sigma-TE	2.0	2.0
Sigma-SA	0.4	0.4
Score pictogram		
Regression line	$0.0 + 0.858 \cdot x$	$0.0 + 0.858 \cdot x$

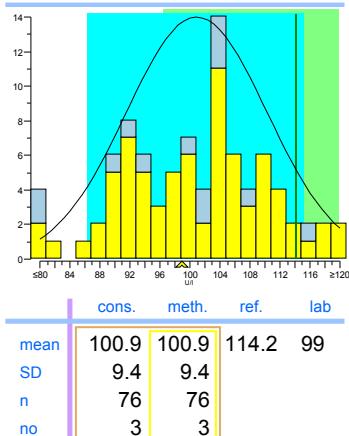
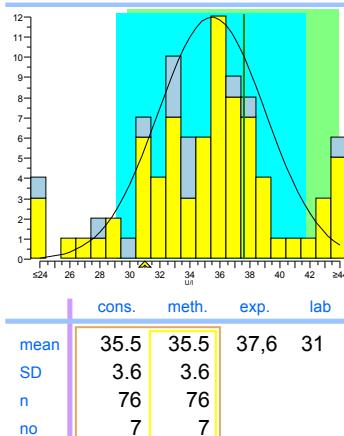
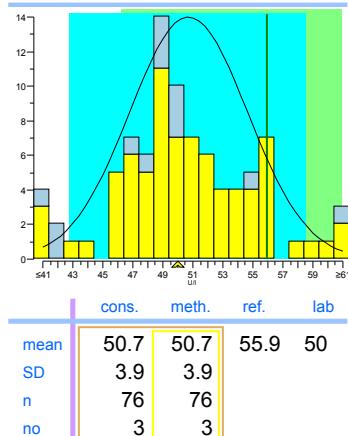
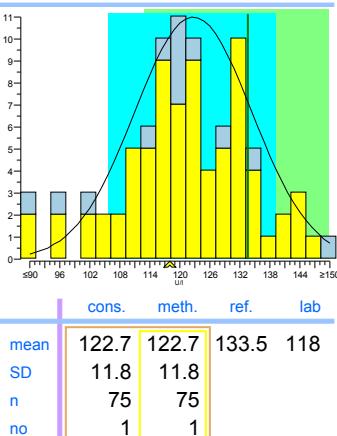
Consensus group IFCC traceerbaar  
 Method IFCC traceable

2013.1 A

2013.1 B

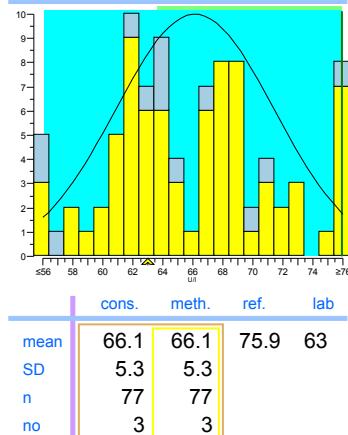
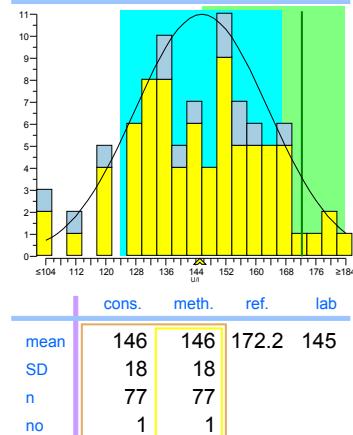
2013.1 C

2013.1 D



2013.1 E

2013.1 F



Legend

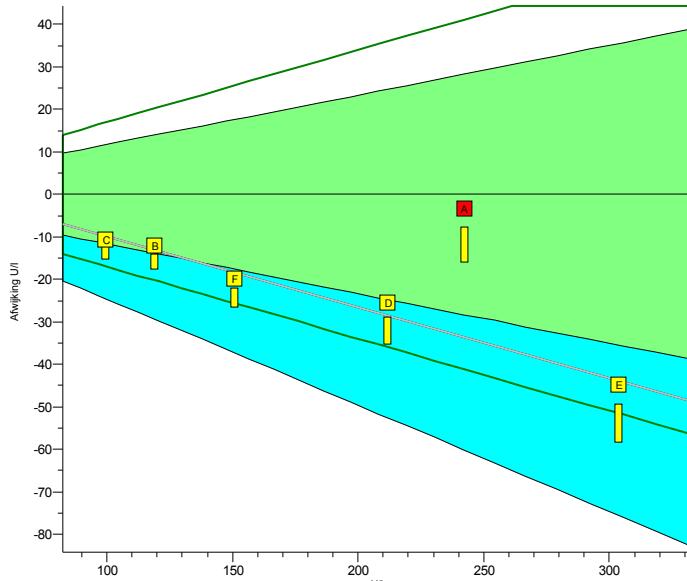
IFCC traceable

IFCC non-traceable

## INPUTs 2013.1

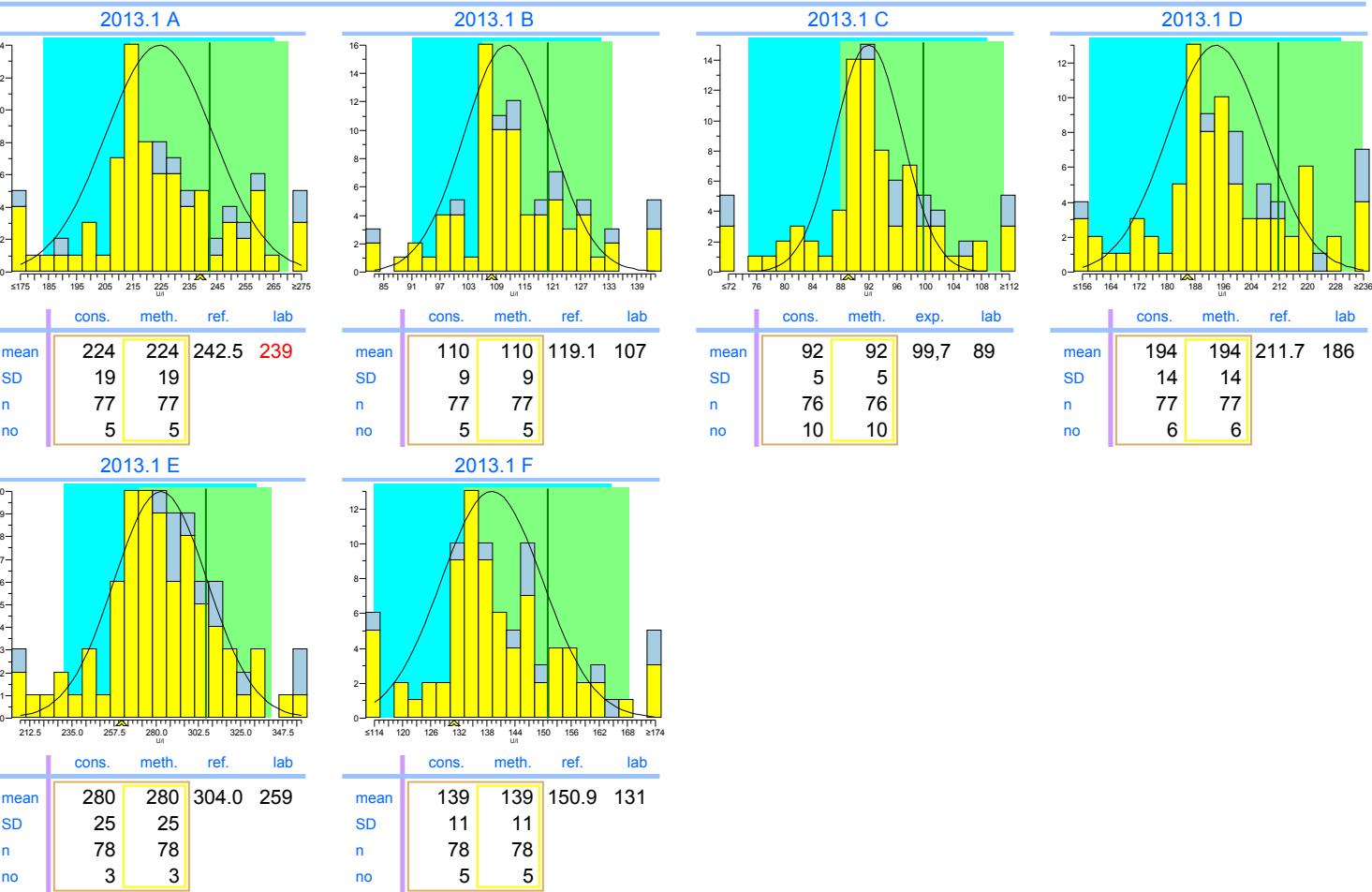
Alk. Phosphatase

units: U/l



	2013.1	cumulative
Trueness	-13%	-13%
Precision	1.3%	1.3%
Number	6	6
Outliers	1	1
Sigma-TE	1.7	1.7
Sigma-SA	5.1 [2]	5.1 [2]
Score pictogram	[Green]	[Green]
Regression line	$7 + 0.834 \cdot x$	$7 + 0.834 \cdot x$

Consensus group IFCC traceerbaar  
 Method IFCC traceable



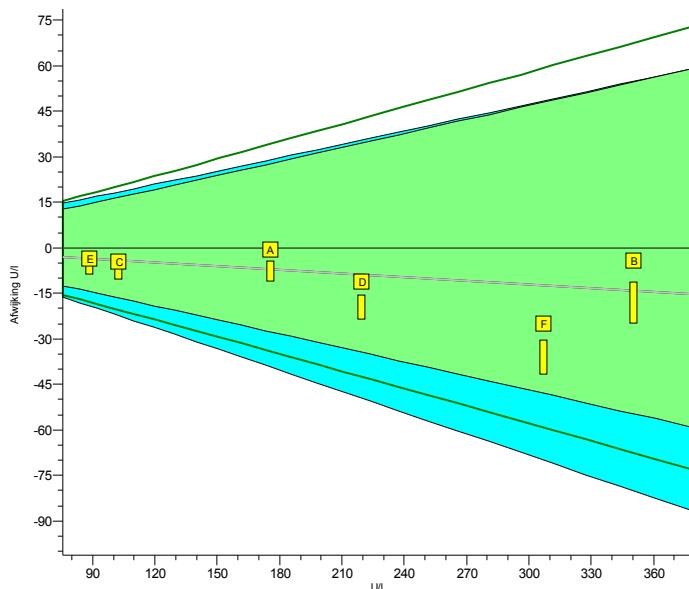
Legend

[Yellow] IFCC traceable      [Grey] IFCC non-traceable

## INPUTs 2013.1

### Amylase

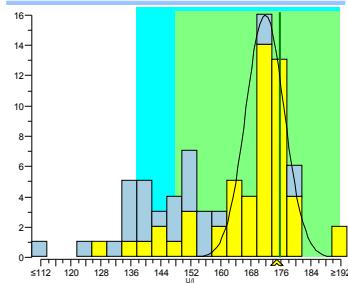
units: U/l



	2013.1	cumulative
Trueness	-4.0%	-4.0%
Precision	3.7%	3.7%
Number	6	6
Outliers	0	0
Sigma-TE	6.0	6.0
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow]	[Yellow]
Regression line	$0 + 0.960 \cdot x$	$0 + 0.960 \cdot x$

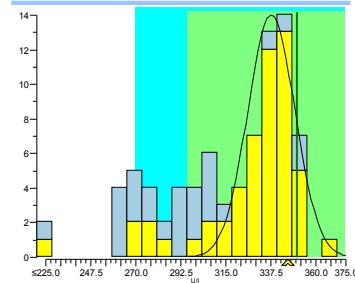
Consensus group  
 Method

2013.1 A



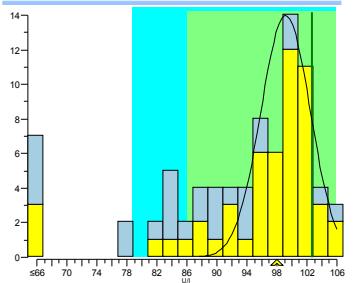
	cons.	meth.	ref.	lab
mean	172	172	175.8	175
SD	5	5		
n	53	53		
no	11	11		

2013.1 B



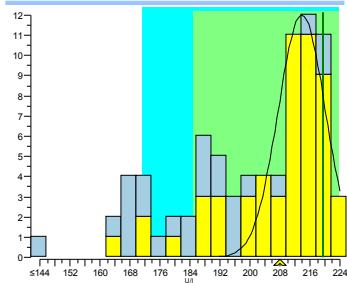
	cons.	meth.	ref.	lab
mean	338	338	350.5	346
SD	12	12		
n	53	53		
no	7	7		

2013.1 C



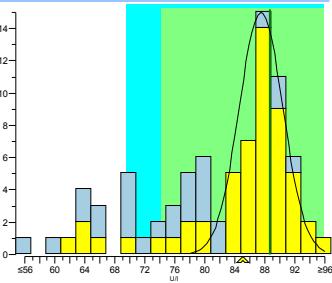
	cons.	meth.	exp.	lab
mean	99	99	102,7	98
SD	3	3		
n	53	53		
no	8	8		

2013.1 D



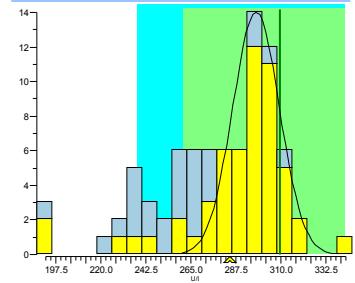
	cons.	meth.	ref.	lab
mean	214	214	219,4	208
SD	6	6		
n	54	54		
no	10	10		

2013.1 E



	cons.	meth.	ref.	lab
mean	87	87	88,7	85
SD	3	3		
n	54	54		
no	10	10		

2013.1 F



	cons.	meth.	ref.	lab
mean	295	295	307,0	282
SD	12	12		
n	54	54		
no	8	8		

## Legend

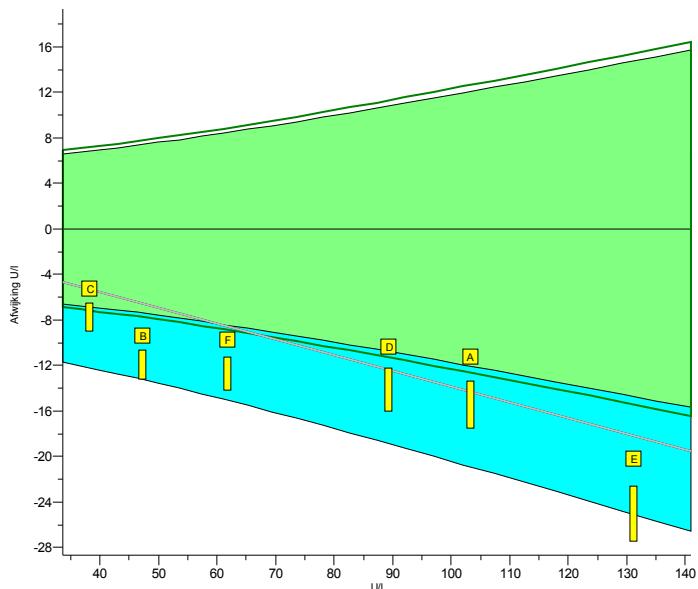
■ IFCC traceable

■ IFCC non-traceable

## INPUTS 2013.1

ASAT

units: U/l

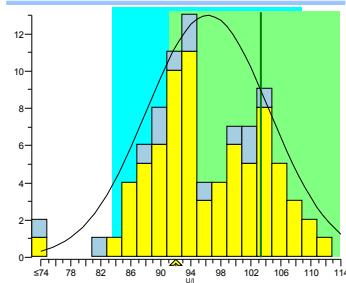


	2013.1	cumulative
Trueness	-14%	-14%
Precision	2.8%	2.8%
Number	6	6
Outliers	0	0
Sigma-TE	0.3	0.3
Sigma-SA	0.6	0.6
Score pictogram		
Regression line	$0.0 + 0.862 \cdot x$	$0.0 + 0.862 \cdot x$

Consensus group  
 Method

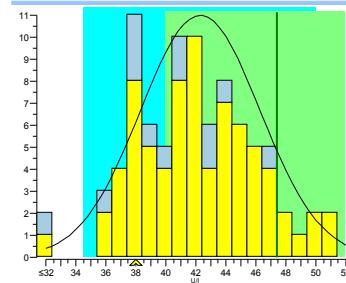
IFCC traceerbaar  
 IFCC traceable

2013.1 A



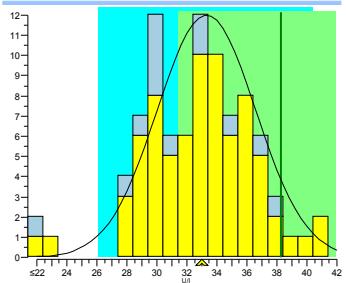
	cons.	meth.	ref.	lab
mean	96.3	96.3	103.3	92
SD	8.2	8.2		
n	75	75		
no	1	1		

2013.1 B



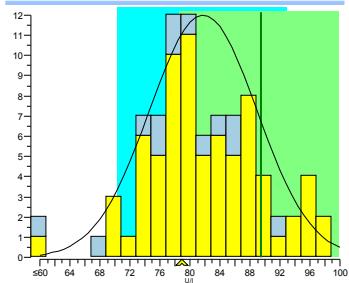
	cons.	meth.	ref.	lab
mean	42.3	42.3	47.4	38
SD	4.0	4.0		
n	75	75		
no	1	1		

2013.1 C



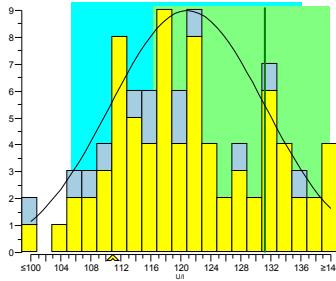
	cons.	meth.	exp.	lab
mean	33.3	33.3	38,3	33
SD	3.3	3.3		
n	75	75		
no	2	2		

2013.1 D



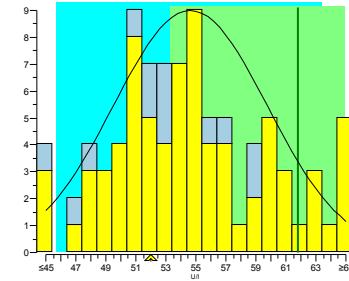
	cons.	meth.	ref.	lab
mean	81.8	81.8	89.4	79
SD	7.2	7.2		
n	74	74		
no	1	1		

2013.1 E



	cons.	meth.	ref.	lab
mean	120.9	120.9	131.2	111
SD	10.3	10.3		
n	76	76		
no	1	1		

2013.1 F



	cons.	meth.	ref.	lab
mean	54.6	54.6	61.8	52
SD	5.1	5.1		
n	76	76		
no	2	2		

Legend

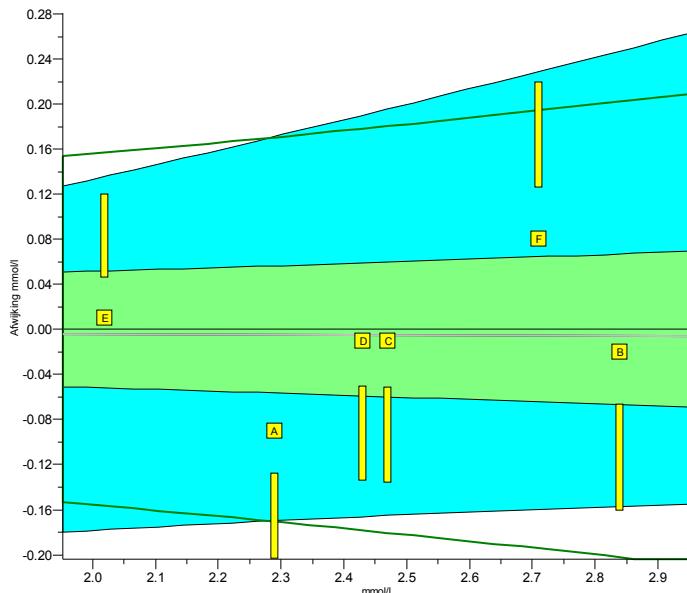
IFCC traceable

IFCC non-traceable

## INPUTs 2013.1

Calcium

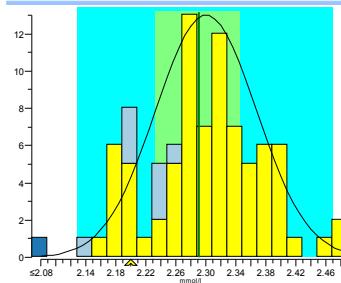
units: mmol/l



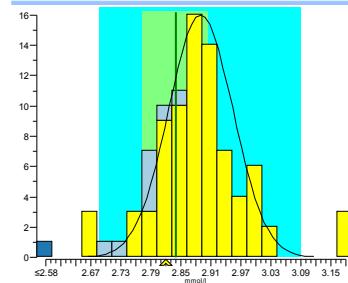
	2013.1	cumulative
Trueness	-0.27%	-0.27%
Precision	2.3%	2.3%
Number	6	6
Outliers	0	0
Sigma-TE	1.5	1.5
Sigma-SA	5.0 [2]	5.0 [2]
Score pictogram	[Yellow]	[Yellow]
Regression line	$0.00 + 0.998 \cdot x$	$0.00 + 0.998 \cdot x$

Consensus group: Colorimetrisch  
 Method: Colorimetric, automatic, discrete

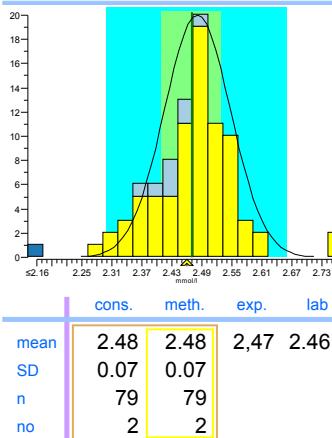
2013.1 A



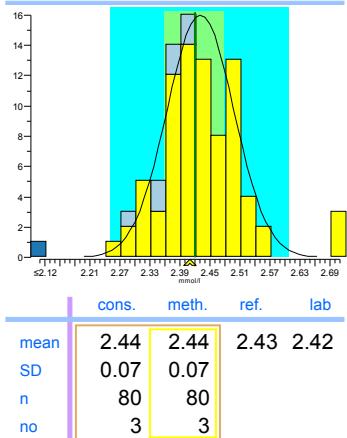
2013.1 B



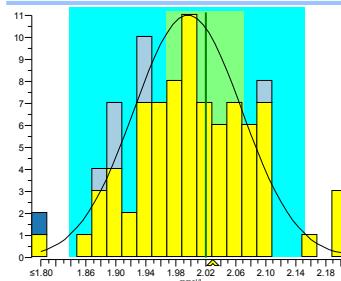
2013.1 C



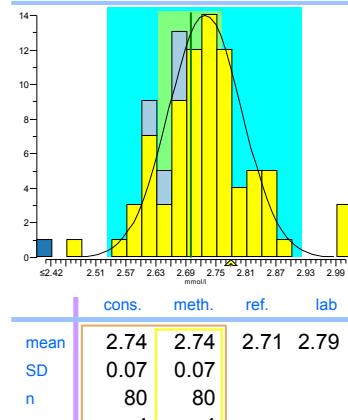
2013.1 D



2013.1 E



2013.1 F



Legend

[Yellow] Colorimetric, automatic, discrete

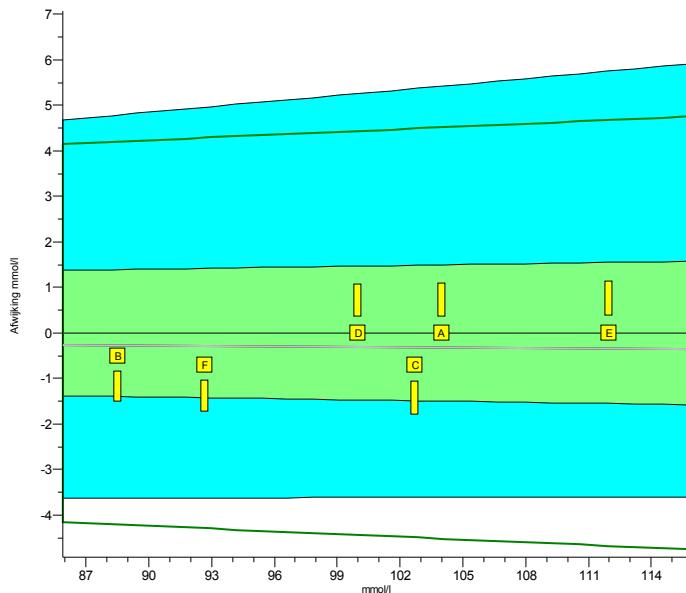
[Grey] ISE indirect (with predilution)

[Blue] Other methods

## INPUTS 2013.1

### Chloride

units: mmol/l



	2013.1	cumulative
Trueness	-0.32%	-0.32%
Precision	0.29%	0.29%
Number	6	6
Outliers	0	0
Sigma-TE	4.7	4.7
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram		
Regression line	$0.0 + 0.997 \cdot x$	$0.0 + 0.997 \cdot x$

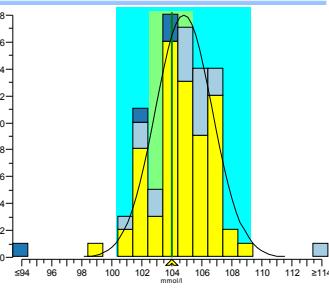
Consensus group ISE/Colorimetric  
 Method ISE indirect (with predilution)

2013.1 A

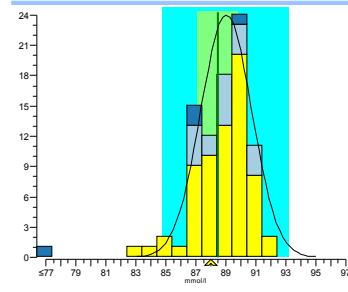
2013.1 B

2013.1 C

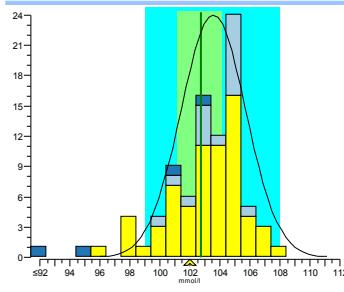
2013.1 D



	cons.	meth.	ref.	lab
mean	104.8	104.8	104	104
SD	1.9	1.8		
n	88	67		
no	3	1		



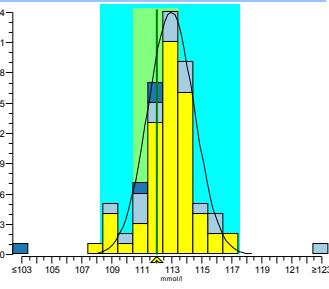
	cons.	meth.	ref.	lab
mean	89.0	89.0	88.5	88
SD	1.7	1.7		
n	88	67		
no	2	1		



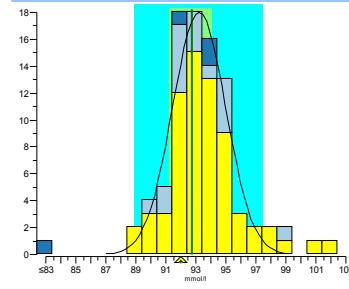
	cons.	meth.	exp.	lab
mean	103.6	103.3	102,7	102
SD	2.2	2.3		
n	88	67		
no	3	1		

2013.1 E

2013.1 F



	cons.	meth.	ref.	lab
mean	113.0	113.0	112	112
SD	1.5	1.7		
n	88	67		
no	3	1		



	cons.	meth.	ref.	lab
mean	93.2	93.3	92.7	92
SD	1.8	1.9		
n	88	67		
no	5	3		

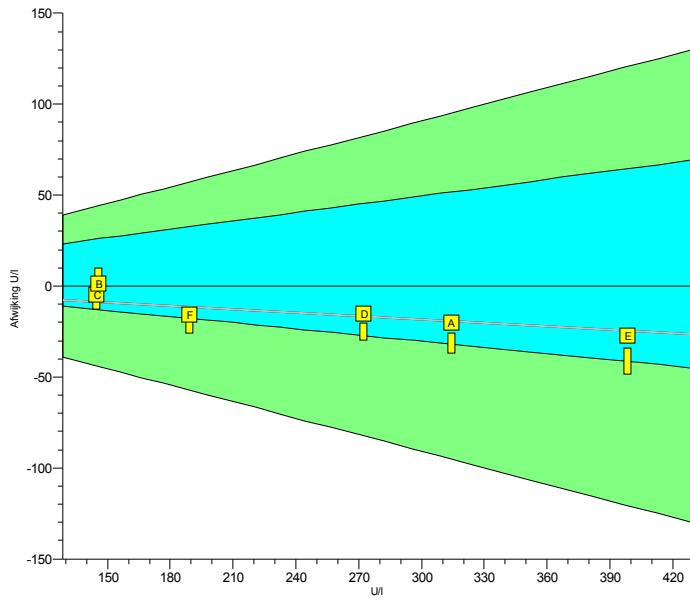
## Legend

ISE indirect (with predilution)	ISE direct (no predilution)	Other methods
---------------------------------	-----------------------------	---------------

## INPUTS 2013.1

CK

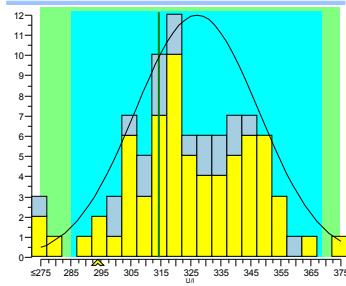
units: U/l



	2013.1	cumulative
Trueness	-5.7%	-5.7%
Precision	1.7%	1.7%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	5.9	5.9
Score pictogram	[Green box with yellow dot]	[Green box with yellow dot]
Regression line	0 + 0.939.x	0 + 0.939.x

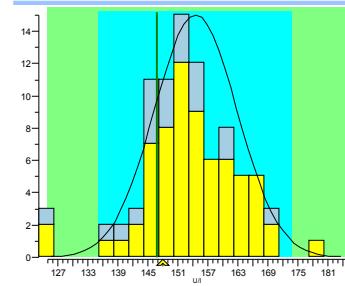
Consensus group IFCC traceerbaar  
 Method IFCC traceable

2013.1 A



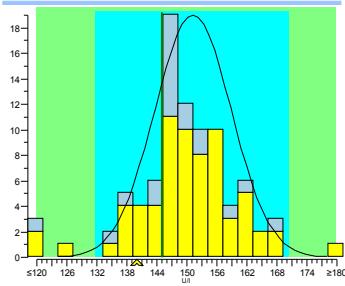
	cons.	meth.	ref.	lab
mean	327	327	314.3	294
SD	21	21		
n	68	68		
no	2	2		

2013.1 B



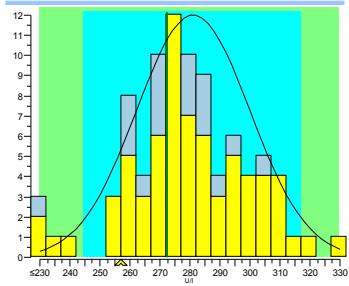
	cons.	meth.	ref.	lab
mean	154	154	145.8	147
SD	8	8		
n	67	67		
no	3	3		

2013.1 C



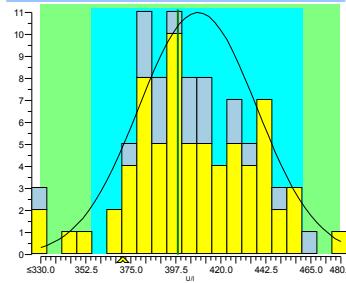
	cons.	meth.	exp.	lab
mean	151	151	145	140
SD	8	8		
n	68	68		
no	4	4		

2013.1 D



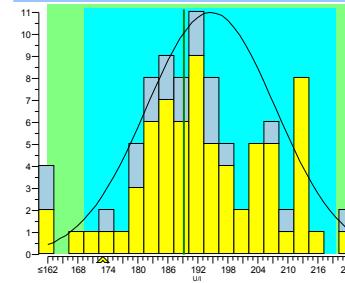
	cons.	meth.	ref.	lab
mean	281	281	272.3	257
SD	19	19		
n	69	69		
no	2	2		

2013.1 E



	cons.	meth.	ref.	lab
mean	409	409	398.4	371
SD	30	30		
n	69	69		
no	2	2		

2013.1 F



	cons.	meth.	ref.	lab
mean	195	195	189.2	173
SD	13	13		
n	69	69		
no	2	2		

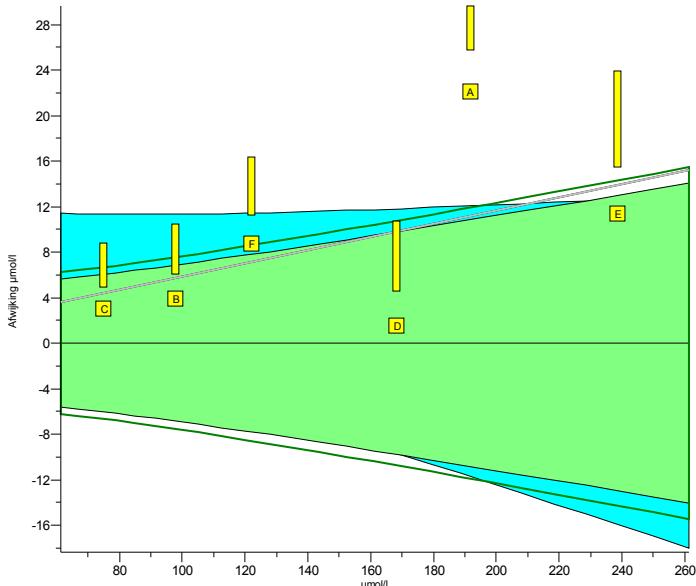
## Legend

[Yellow box] IFCC traceable

[Grey box] IFCC non-traceable

## INPUTs 2013.1

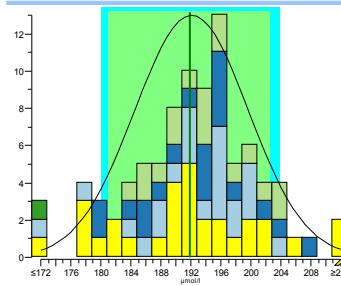
## Creatinine

units:  $\mu\text{mol/l}$ 

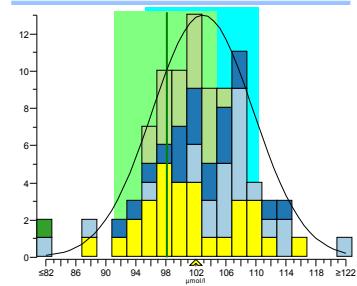
	2013.1	cumulative
Trueness	+5.6%	+5.6%
Precision	4.9%	4.9%
Number	6	6
Outliers	0	0
Sigma-TE	1.4	1.4
Sigma-SA	1.7	1.7
Score pictogram		
Regression line	$0.0 + 1.058.x$	$0.0 + 1.058.x$

Consensus group: Jaffe  
 Method: Alk. Picrate, kinetic with compensation

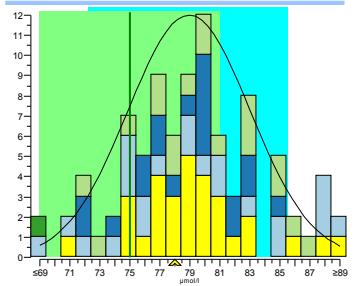
2013.1 A



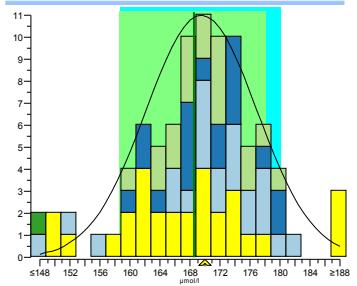
2013.1 B



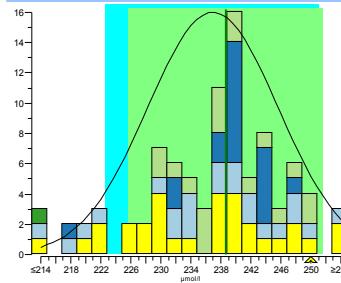
2013.1 C



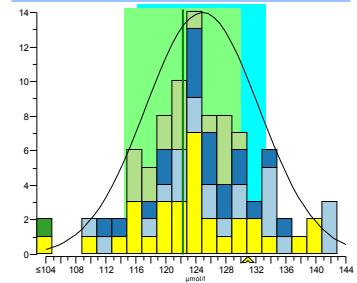
2013.1 D



2013.1 E

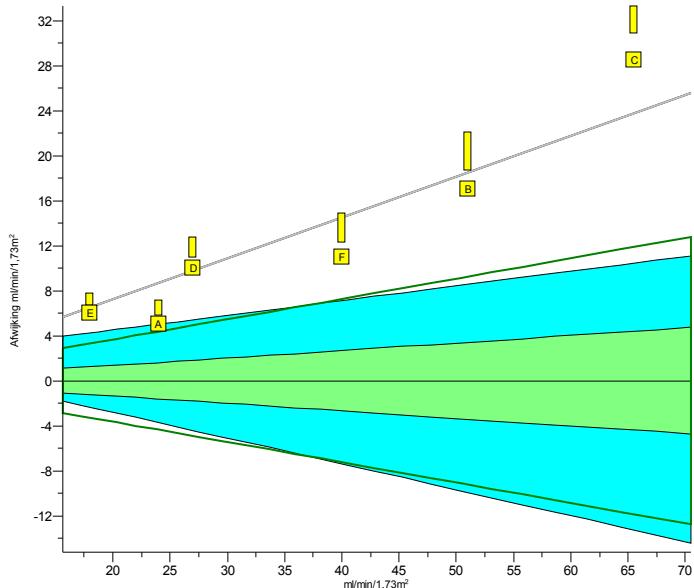


2013.1 F



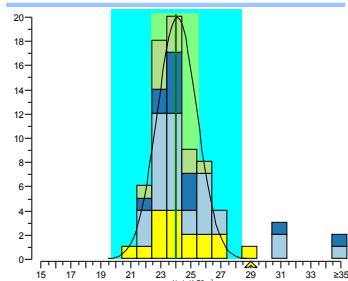
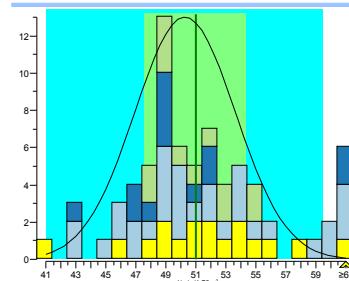
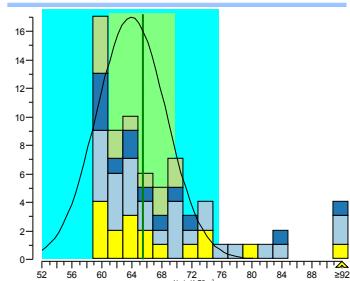
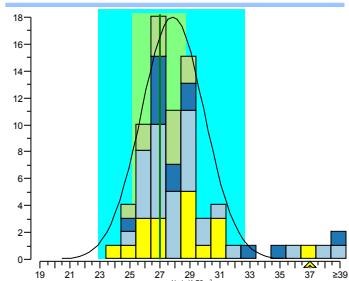
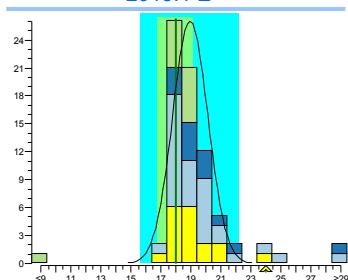
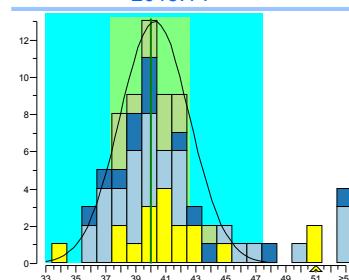
## Legend

- █ Alk. Picrate, kinetic with compensation
- █ Alk. Picrate, kinetic
- █ Alk. Picrate, endpoint
- █ Enzymatic, automatic
- █ Other methods

**INPUTs 2013.1****eGFR (F, 55, white)**units: ml/min/1,73m<sup>2</sup>

	2013.1	cumulative
Trueness	+34%	+34%
Precision	7.0%	7.0%
Number	6	6
Outliers	0	0
Sigma-TE	-3.0	-3.0
Sigma-SA	-2.6 <span style="color:red">0</span>	-2.6 <span style="color:red">0</span>
Score pictogram		
Regression line	$0.0 + 1.363 \cdot x$	$0.0 + 1.363 \cdot x$

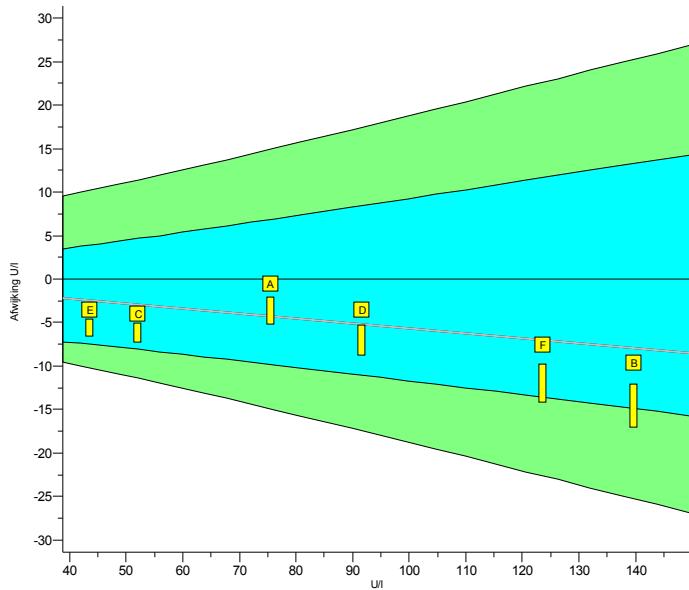
Consensus group: Jaffe  
 Method: Alk. Picrate, kinetic with compensation

**2013.1 A****2013.1 B****2013.1 C****2013.1 D****2013.1 E****2013.1 F****Legend**■ Alk. Picrate, kinetic with compensation■ Other methods■ Alk. Picrate, kinetic■ Enzymatic, automatic

**INPUTs 2013.1**

Gamma-GT

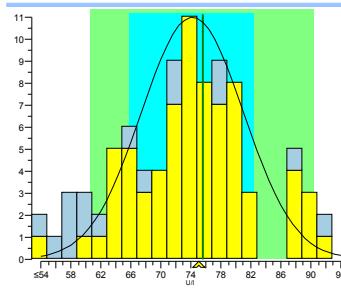
units: U/l



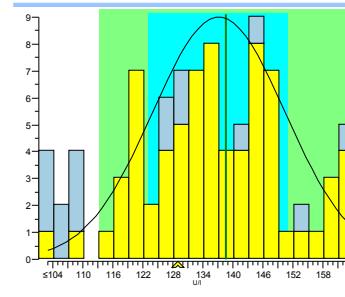
	2013.1	cumulative
Trueness	-5.5%	-5.5%
Precision	2.7%	2.7%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	4.1	4.1
Score pictogram	[Green box]	[Green box]
Regression line	$0.0 + 0.943 \cdot x$	$0.0 + 0.943 \cdot x$

 Consensus group: IFCC traceerbaar  
 Method: IFCC traceable

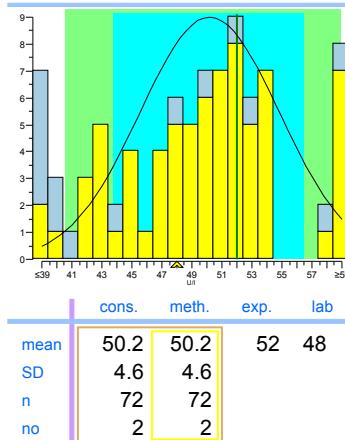
2013.1 A



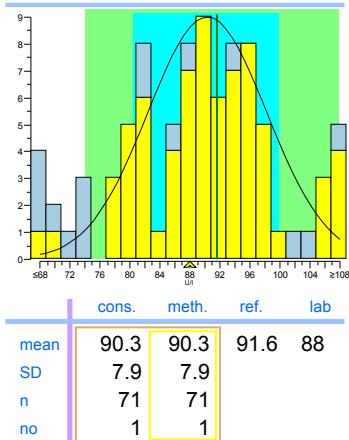
2013.1 B



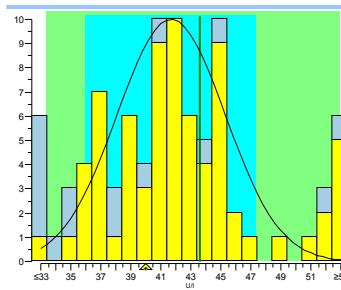
2013.1 C



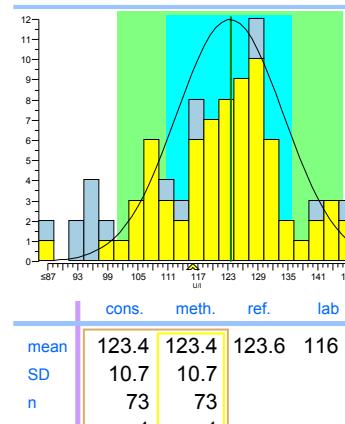
2013.1 D



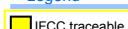
2013.1 E

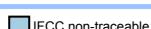


2013.1 F



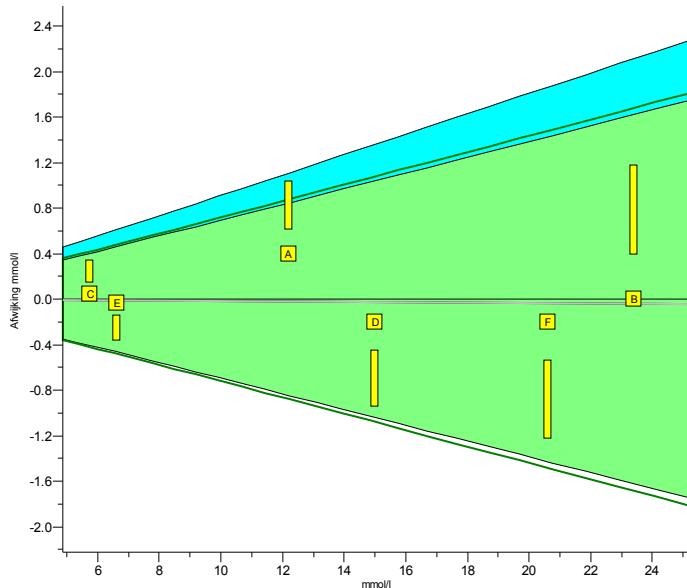
## Legend

 IFCC traceable

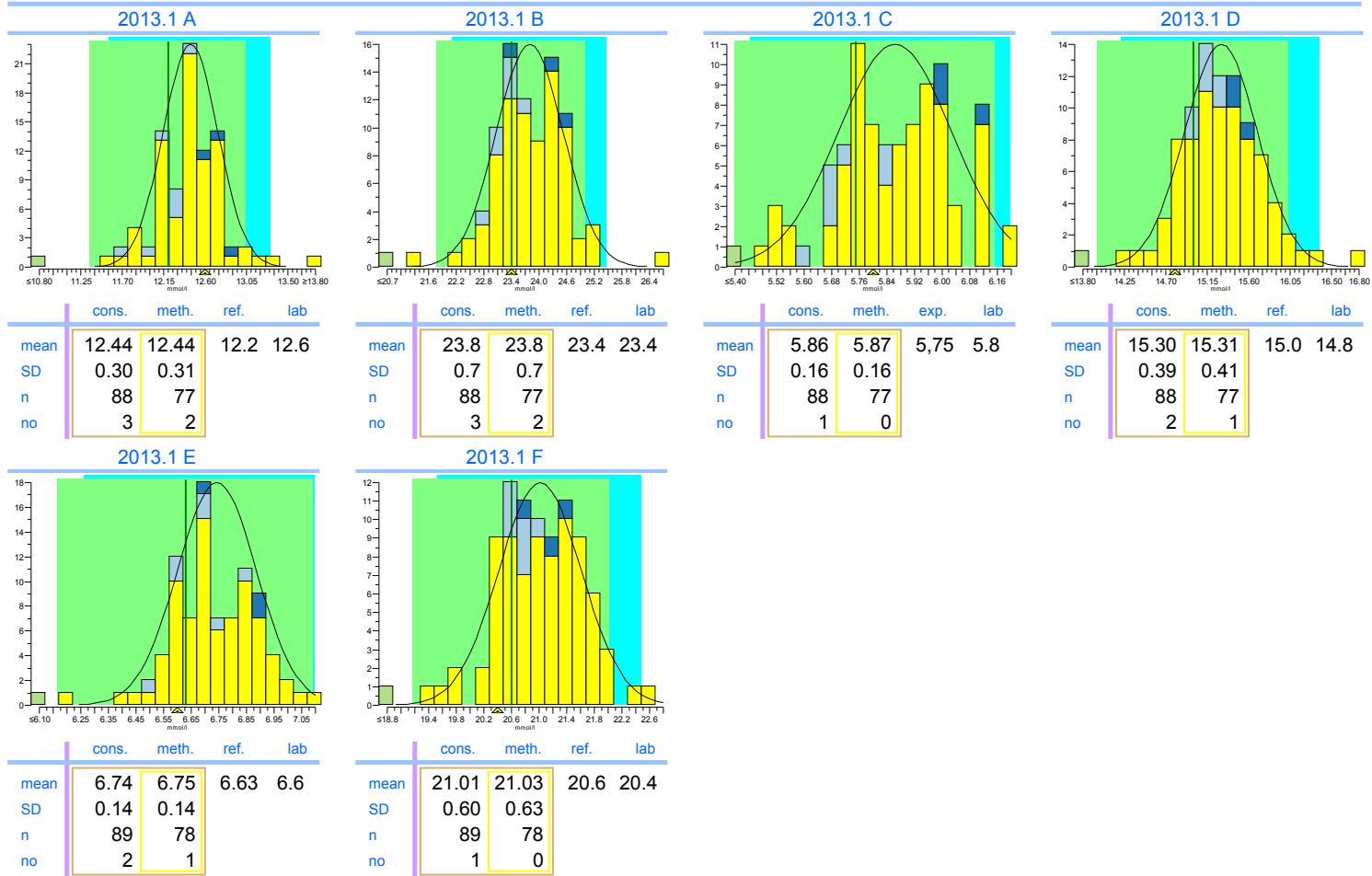
 IFCC non-traceable

**INPUTs 2013.1****Glucose**

units: mmol/l



	2013.1	cumulative
Trueness	+0.02%	+0.02%
Precision	1.6%	1.6%
Number	6	6
Outliers	0	0
Sigma-TE	5.0	5.0
Sigma-SA	5.2 [2]	5.2 [2]
Score pictogram	[Green]	[Green]
Regression line	$0.00 + 0.998 \cdot x$	$0.00 + 0.998 \cdot x$

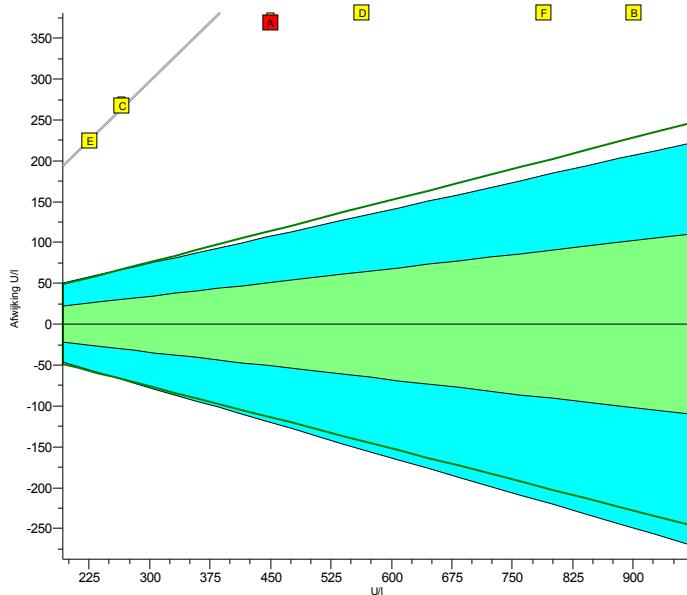
 Consensus group: Natte chemie  
 Method: Hexokinase, automatic
**Legend**

- [Yellow] Hexokinase, automatic
- [Light Blue] Glucose-oxydase, amperometric, H<sub>2</sub>O<sub>2</sub>
- [Dark Blue] Glucose-oxydase/POD,automatic
- [Green] Other methods

## INPUTs 2013.1

LD

units: U/l

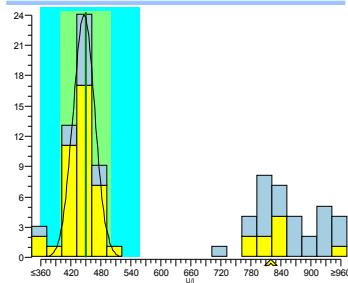


	2013.1	cumulative
Trueness	+98%	+98%
Precision	1.1%	1.1%
Number	6	6
Outliers	1	1
Sigma-TE	-3.0	-3.0
Sigma-SA	-3.0	-3.0
Score pictogram	0	0
Regression line	$10 + 1.958 \cdot x$	$10 + 1.958 \cdot x$

Consensus group  
Method

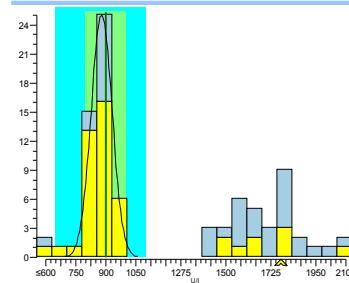
IFCC traceerbaar  
IFCC traceable

2013.1 A



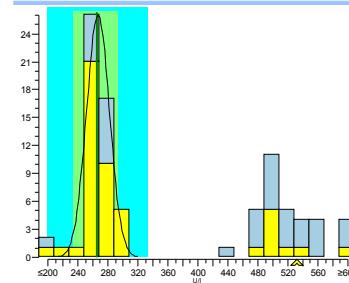
	cons.	meth.	ref.	lab
mean	446	446	450.7	820
SD	22	22		
n	48	48		
no	11	11		

2013.1 B



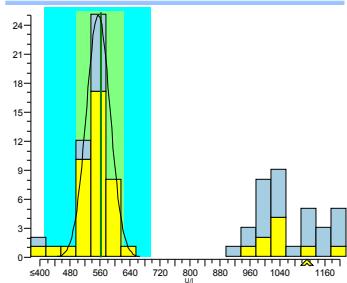
	cons.	meth.	ref.	lab
mean	879	879	900.8	1773
SD	52	52		
n	47	47		
no	11	11		

2013.1 C



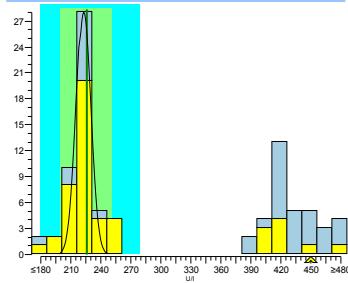
	cons.	meth.	exp.	lab
mean	267	267	265	532
SD	15	15		
n	48	48		
no	11	11		

2013.1 D



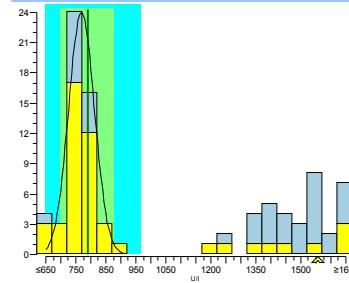
	cons.	meth.	ref.	lab
mean	556	556	563.0	1111
SD	32	32		
n	48	48		
no	11	11		

2013.1 E



	cons.	meth.	ref.	lab
mean	223	223	226.1	450
SD	7	7		
n	48	48		
no	16	16		

2013.1 F



	cons.	meth.	ref.	lab
mean	769	769	788.9	1556
SD	43	43		
n	48	48		
no	11	11		

## Legend

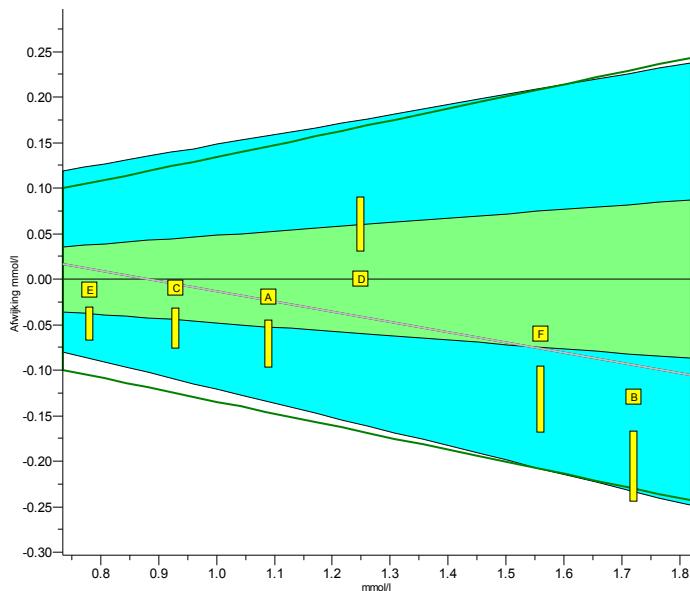
 IFCC traceable

 IFCC non-traceable

## INPUTs 2013.1

### Magnesium

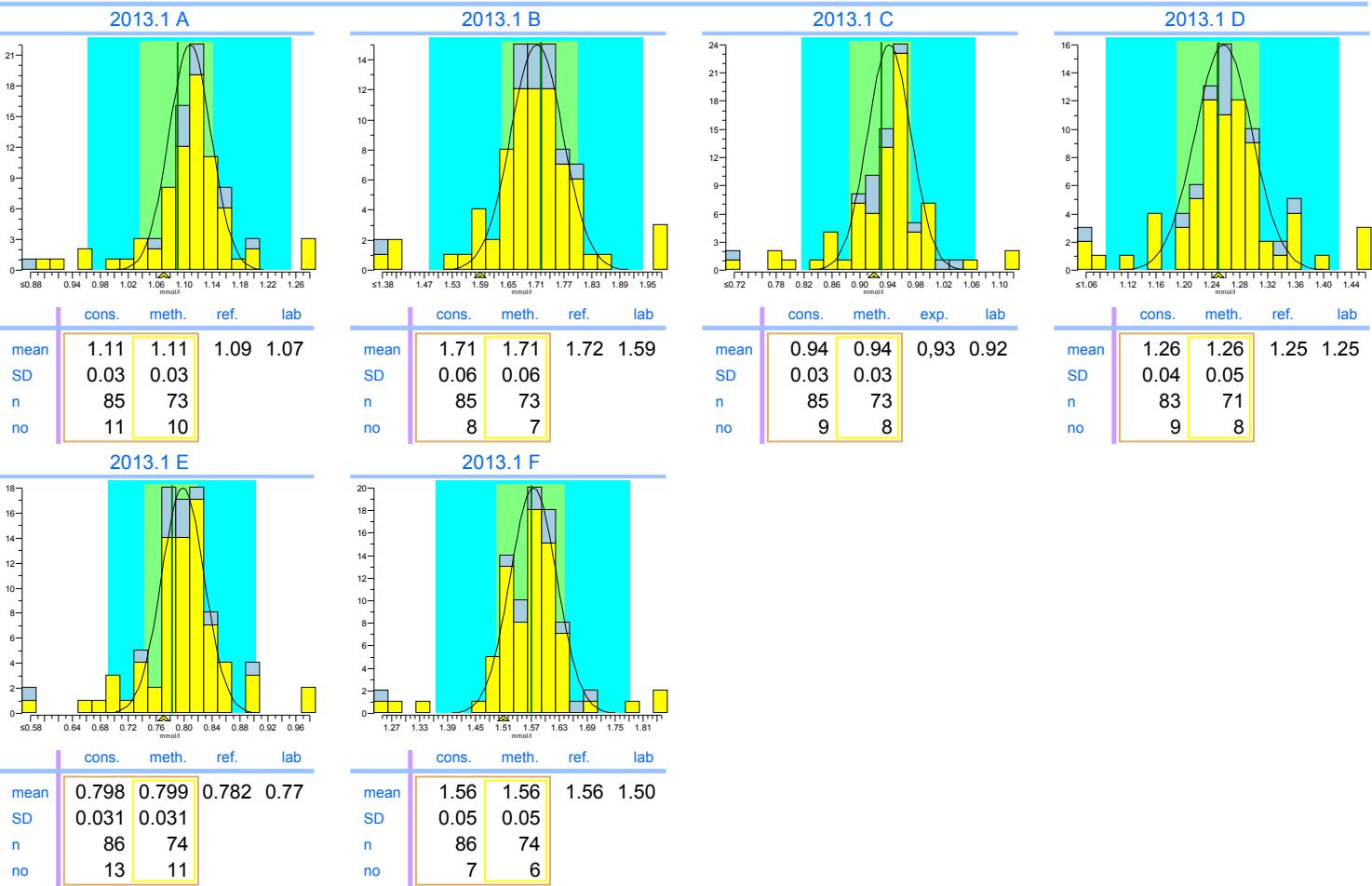
units: mmol/l



	2013.1	cumulative
Trueness	-3.2%	-3.2%
Precision	2.6%	2.6%
Number	6	6
Outliers	0	0
Sigma-TE	1.9	1.9
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Score pictogram icon]	[Score pictogram icon]
Regression line	$0.10 + 0.887 \cdot x$	$0.10 + 0.887 \cdot x$

Consensus group  
Method

Overall  
Colorimetric

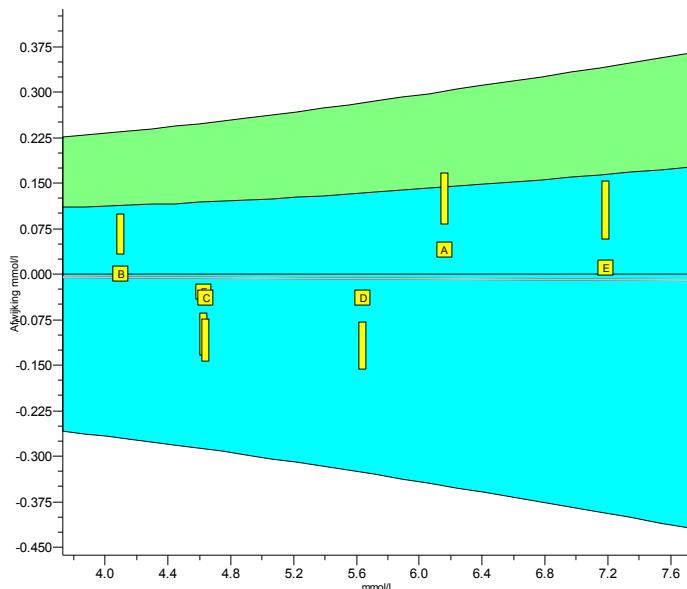


#### Legend



**INPUTs 2013.1****Potassium**

units: mmol/l



	2013.1	cumulative
Trueness	-0.19%	-0.19%
Precision	0.59%	0.59%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	6.0	6.0
Score pictogram	[Color-coded box]	[Color-coded box]
Regression line	$0.00 + 0.999 \cdot x$	$0.00 + 0.999 \cdot x$

 Consensus group ISE verdund/Vlamfotometrie  
 Method ISE indirect (with predilution)

2013.1 A

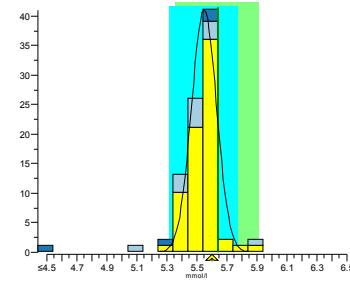
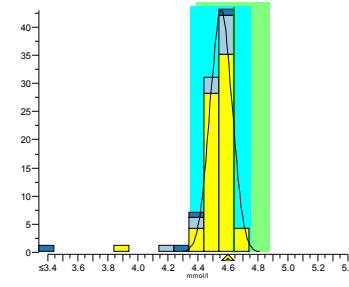
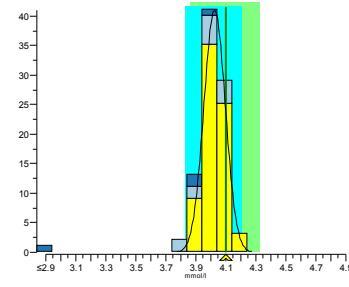
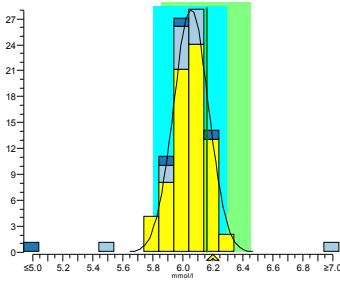
2013.1 B

2013.1 C

2013.1 D

2013.1 E

2013.1 F

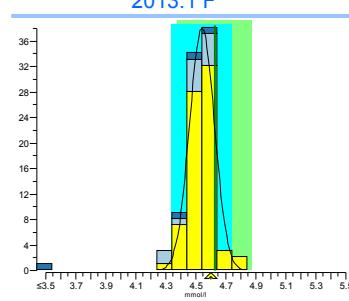
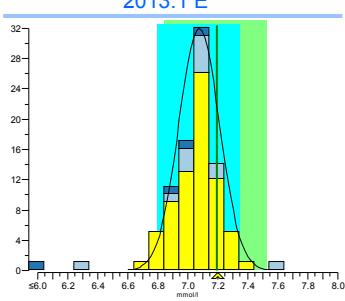


	cons.	meth.	ref.	lab
mean	6.05	6.05	6.16	6.2
SD	0.12	0.12		
n	72	72		
no	0	0		

	cons.	meth.	ref.	lab
mean	4.02	4.02	4.10	4.1
SD	0.07	0.07		
n	72	72		
no	0	0		

	cons.	meth.	exp.	lab
mean	4.55	4.55	4.64	4.6
SD	0.07	0.07		
n	72	72		
no	1	1		

	cons.	meth.	ref.	lab
mean	5.55	5.55	5.64	5.6
SD	0.08	0.08		
n	72	72		
no	2	2		



	cons.	meth.	ref.	lab
mean	7.08	7.08	7.19	7.2
SD	0.14	0.14		
n	72	72		
no	0	0		

	cons.	meth.	ref.	lab
mean	4.54	4.54	4.63	4.6
SD	0.08	0.08		
n	73	73		
no	2	2		

## Legend

ISE indirect (with predilution)

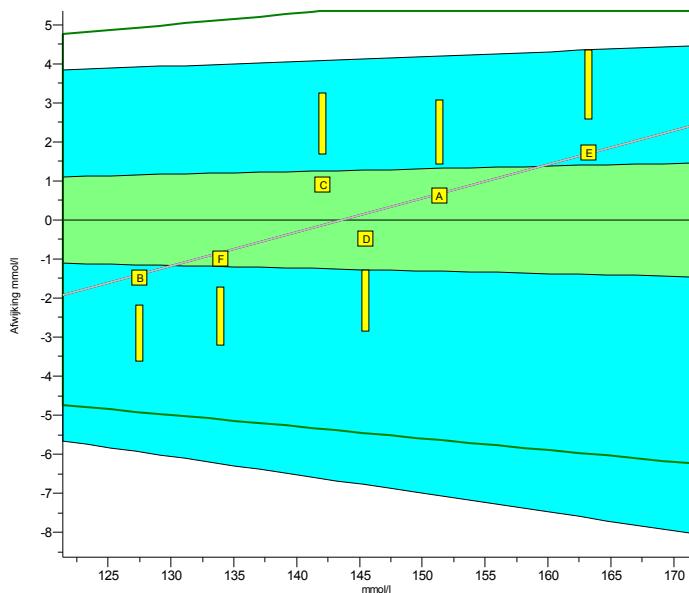
ISE direct (no predilution)

Other methods

## INPUTS 2013.1

Sodium

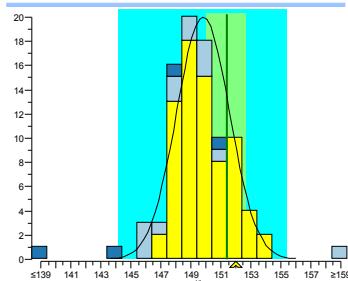
units: mmol/l



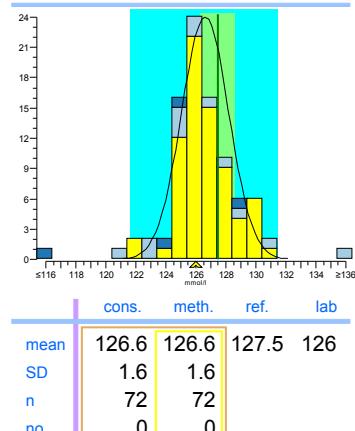
	2013.1	cumulative
Trueness	+0.02%	+0.02%
Precision	0.45%	0.45%
Number	6	6
Outliers	0	0
Sigma-TE	1.2	1.2
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow]	[Yellow]
Regression line	$-12.4 + 1.086 \cdot x$	$-12.4 + 1.086 \cdot x$

 Consensus group ISE verdund/Vlamfotometrie  
 Method ISE indirect (with predilution)

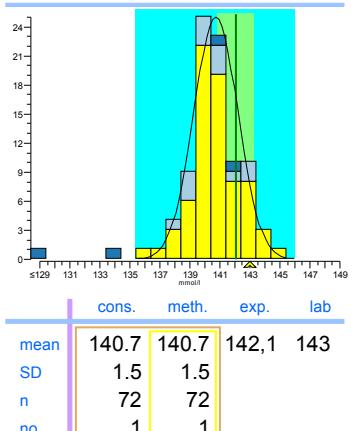
2013.1 A



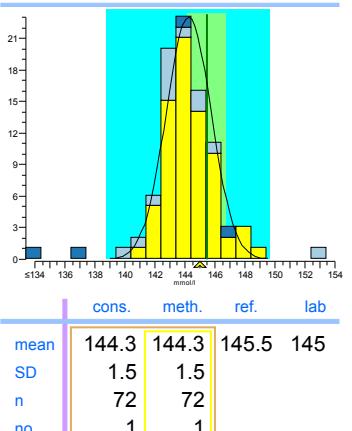
2013.1 B



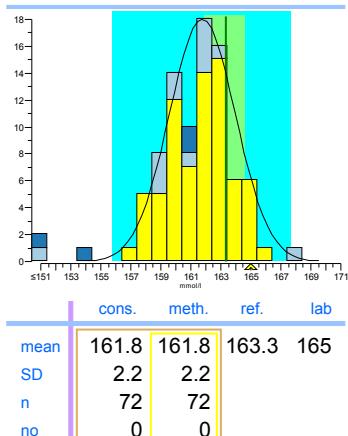
2013.1 C



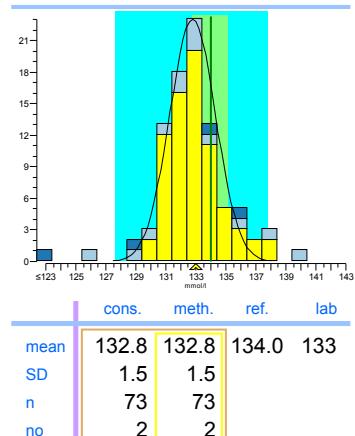
2013.1 D



2013.1 E



2013.1 F



## Legend

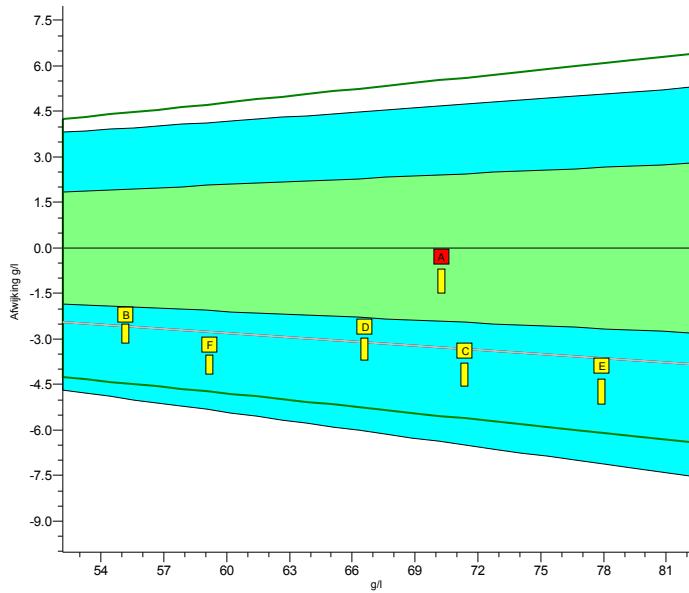
[Yellow] ISE indirect (with predilution)

[Blue] ISE direct (no predilution)

[Grey] Other methods

**INPUTS 2013.1****Total Protein**

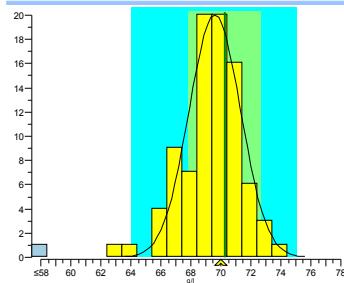
units: g/l



	2013.1	cumulative
Trueness	-4.6%	-4.6%
Precision	0.46%	0.46%
Number	6	6
Outliers	1	1
Sigma-TE	-0.1	-0.1
Sigma-SA	6.0 [2]	6.0 [2]
Score pictogram	[Yellow, Green, Blue]	[Yellow, Green, Blue]
Regression line	$0.0 + 0.953 \cdot x$	$0.0 + 0.953 \cdot x$

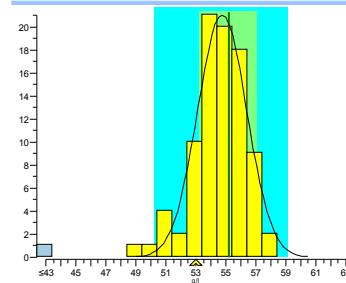
 Consensus group: Biureet  
 Method: Biurete, automatic

2013.1 A



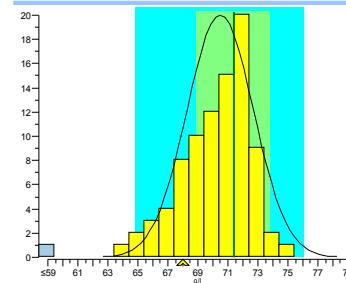
	cons.	meth.	ref.	lab
mean	69.6	69.6	70.3	70
SD	1.7	1.7		
n	88	88		
no	2	2		

2013.1 B



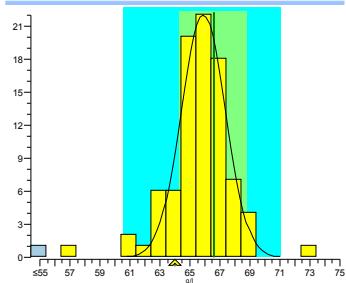
	cons.	meth.	ref.	lab
mean	54.8	54.8	55.2	53
SD	1.6	1.6		
n	88	88		
no	1	1		

2013.1 C



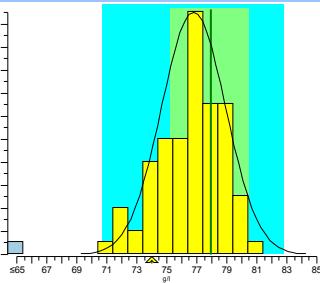
	cons.	meth.	exp.	lab
mean	70.5	70.5	71.4	68
SD	2.2	2.2		
n	87	87		
no	0	0		

2013.1 D



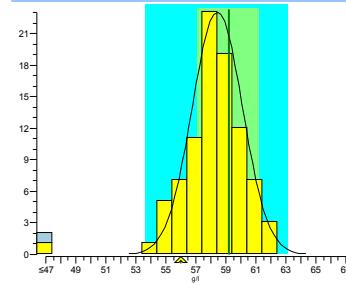
	cons.	meth.	ref.	lab
mean	65.9	65.9	66.6	64
SD	1.5	1.5		
n	88	88		
no	4	4		

2013.1 E



	cons.	meth.	ref.	lab
mean	76.8	76.8	77.9	74
SD	2.1	2.1		
n	88	88		
no	0	0		

2013.1 F



	cons.	meth.	ref.	lab
mean	58.4	58.4	59.2	56
SD	1.7	1.7		
n	89	89		
no	1	1		

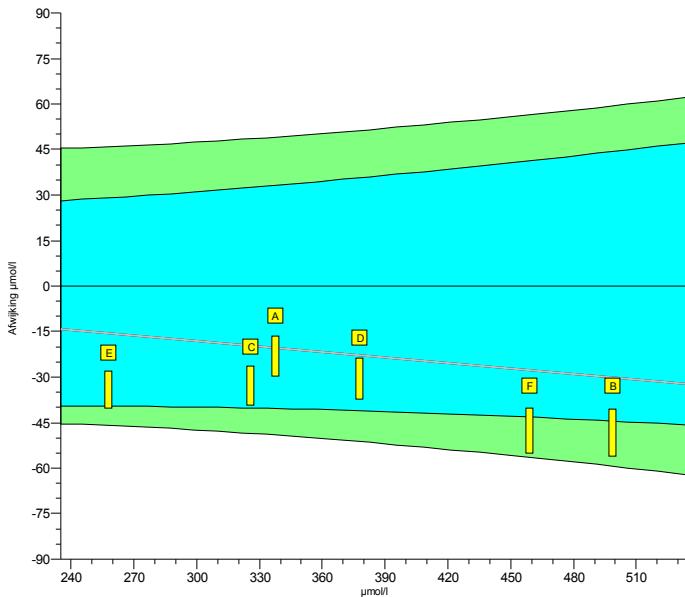
## Legend

 Biurete, automatic

 Other methods

## INPUTs 2013.1

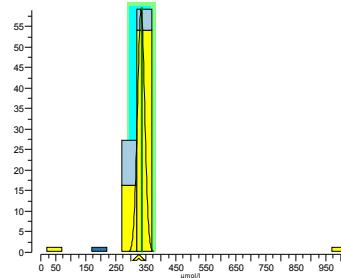
Urate

units:  $\mu\text{mol/l}$ 

	2013.1	cumulative
Trueness	-6.0%	-6.0%
Precision	2.0%	2.0%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	3.7	3.7
Score pictogram	[Yellow box]	[Yellow box]
Regression line	$0 + 0.940 \cdot x$	$0 + 0.940 \cdot x$

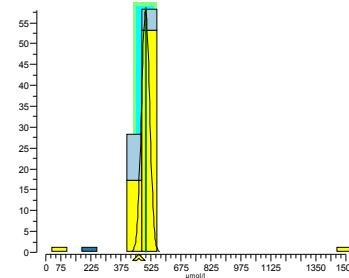
Consensus group: Colorimetrisch  
 Method: Uricase, colorim., automatic

2013.1 A



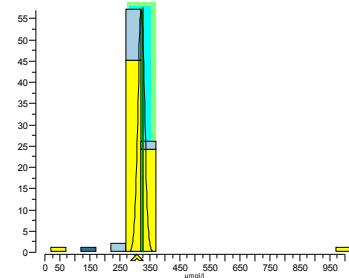
	cons.	meth.	ref.	lab
mean	334	334	0.338	328
SD	12	12		
n	88	72		
no	18	2		

2013.1 B



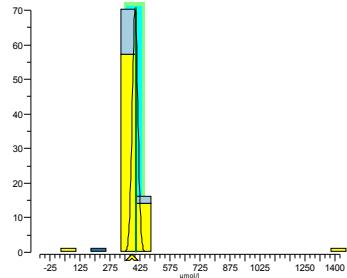
	cons.	meth.	ref.	lab
mean	499	499	0.499	466
SD	20	20		
n	88	72		
no	18	2		

2013.1 C



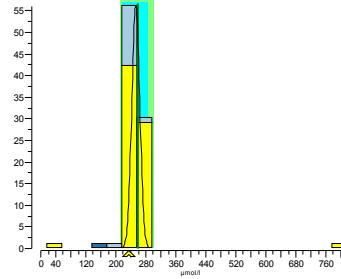
	cons.	meth.	exp.	lab
mean	321	321	0.326	306
SD	11	11		
n	87	71		
no	18	2		

2013.1 D



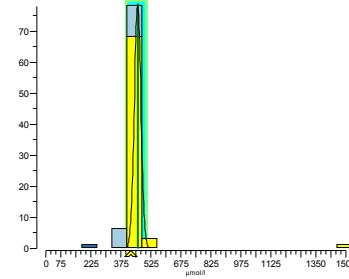
	cons.	meth.	ref.	lab
mean	376	376	0.378	361
SD	14	14		
n	88	73		
no	17	2		

2013.1 E



	cons.	meth.	ref.	lab
mean	253	253	0.258	236
SD	11	11		
n	89	73		
no	18	2		

2013.1 F



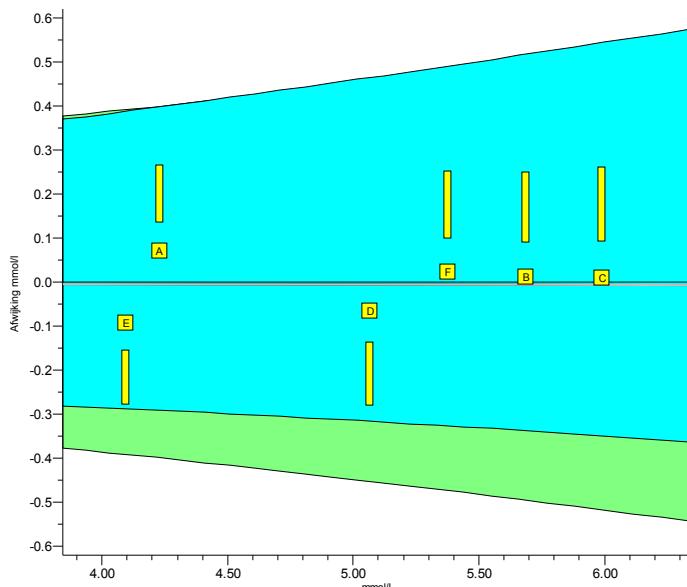
	cons.	meth.	ref.	lab
mean	459	459	0.459	426
SD	17	17		
n	88	72		
no	17	1		

## Legend

[Yellow box] Uricase, colorim., automatic    [Grey box] Uricase, differential UV, automatic    [Blue box] Oudege methoden

**INPUTs 2013.1****Cholesterol**

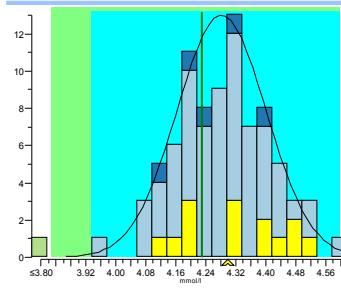
units: mmol/l



	2013.1	cumulative
Trueness	-0.15%	-0.15%
Precision	1.1%	1.1%
Number	6	6
Outliers	0	0
Sigma-TE	6.0 [2]	6.0 [2]
Sigma-SA	6.0	6.0
Score pictogram	[Score pictogram]	[Score pictogram]
Regression line	$0.00 + 0.999 \cdot x$	$0.00 + 0.999 \cdot x$

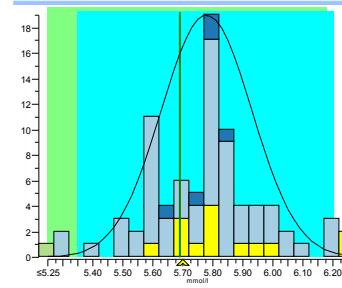
 Consensus group: Enzymatisch  
 Method: Enzymatic, automatic, kinetic

2013.1 A



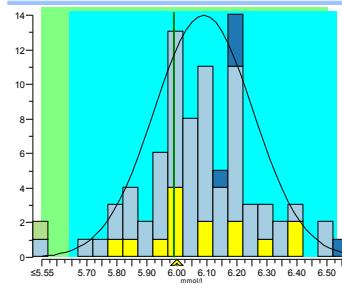
	cons.	meth.	ref.	lab
mean	4.28	4.32	4.230	4.3
SD	0.12	0.13		
n	83	14		
no	0	0		

2013.1 B



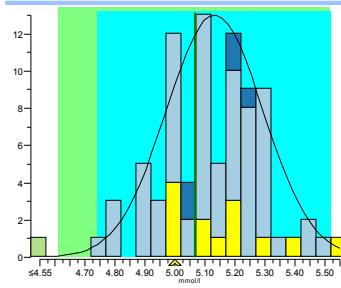
	cons.	meth.	ref.	lab
mean	5.78	5.79	5.689	5.7
SD	0.15	0.12		
n	83	14		
no	4	2		

2013.1 C



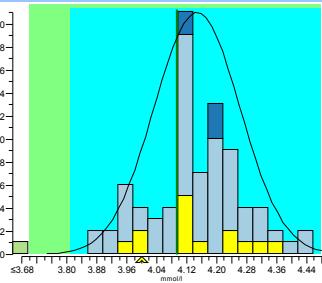
	cons.	meth.	exp.	lab
mean	6.09	6.10	5.99	6.0
SD	0.16	0.19		
n	83	14		
no	1	0		

2013.1 D



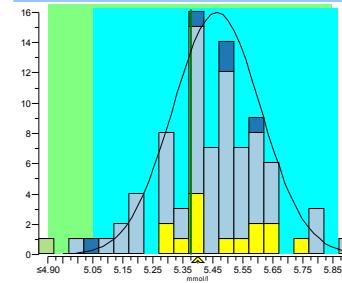
	cons.	meth.	ref.	lab
mean	5.13	5.18	5.066	5.0
SD	0.16	0.18		
n	82	13		
no	0	0		

2013.1 E



	cons.	meth.	ref.	lab
mean	4.15	4.14	4.094	4.0
SD	0.11	0.12		
n	84	14		
no	0	0		

2013.1 F



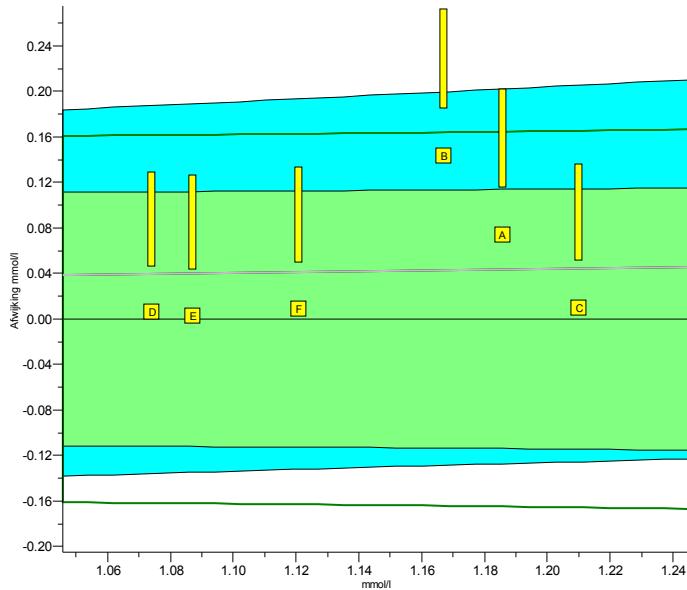
	cons.	meth.	ref.	lab
mean	5.46	5.49	5.377	5.4
SD	0.15	0.14		
n	84	14		
no	2	0		

## Legend

- Enzymatic, automatic, kinetic
- Enzymatic, automatic, discrete
- Abell-Kendall reference values
- Overige methoden

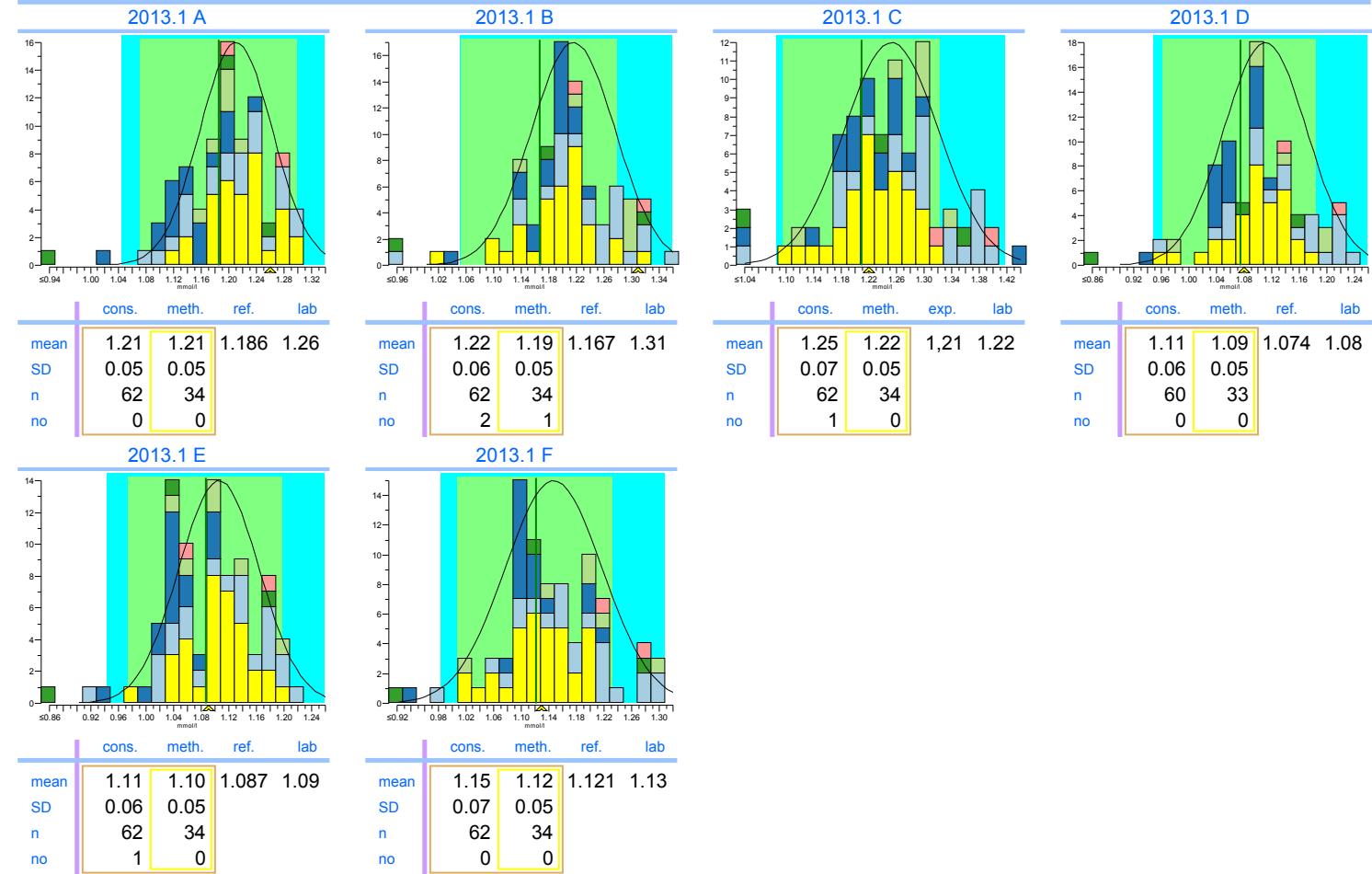
**INPUTS 2013.1****HDL-Cholesterol**

units: mmol/l



	2013.1	cumulative
Trueness	+3.6%	+3.6%
Precision	4.4%	4.4%
Number	6	6
Outliers	0	0
Sigma-TE	2.9	2.9
Sigma-SA	4.4 [1]	4.4 [1]
Score pictogram		
Regression line	$0.00 + 1.037 \cdot x$	$0.00 + 1.037 \cdot x$

Consensus group: Direct  
 Method: PEG modified enzyme, PEGME (Kyowa Medex)

**Legend**

PEG modified enzyme, PEGME (Kyowa Medex)

Accelerator Selective Detergent ("Ultra HDL")

Immunoinhibition

Catalase method (Denka Seiken)

Overige methoden

Precipitation Technique